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177448

From: Mertz, Prema
Sent: Tuesday, January 24, 2006 1:29 PM
To: STIC-Biotech/ChemLib
Subject: 10/767521

Please search SEQ ID NO:1 with protein databases.

Thanks.

Prema Mertz, Ph.D., J.D.
Primary Examiner
Art Unit 1646
4D81 Remsen Bldg Mailbox 4C70
US Patent & Trademark Office
Tel # (571) 272-0876
FAX # (571) 273-0876

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JAN 24, 2006
STIC-BIOTECH/ChemLib
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Searcher: _____
Searcher Phone: _____
Date Searcher Picked up: _____
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Type of Search
NA# _____ AA# _____
S/L: _____ Oligomer: _____
Encode/Transl: _____
Structure #: _____ Text: _____
Inventor: _____ Litigation: _____

Vendors and cost where applicable
STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other (Specify): _____

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Mg

175414

STIC-Biotech/ChemLib

From: Mertz, Prema
Sent: Sunday, January 01, 2006 12:49 PM
To: STIC-Biotech/ChemLib
Subject: 10,767,521

RECEIVED
JAN - 3 - 2006
JH/CIEM/STIC/STIC

Please search SEQ ID NO:2, 3, 4, with DNA databases.

Please search US PG PUB databases with SEQ ID NO:2, 3, and 4.

Thanks.

Prema Mertz, Ph.D., J.D.
Primary Examiner
Art Unit 1646
4D81 Remsen Bldg Mailbox 4C70
US Patent & Trademark Office
Tel # (571) 272-0876
FAX # (571) 273-0876

Searcher: _____
Searcher Phone: _____
Date Searcher Picked up: _____
Date completed: _____
Searcher Prep Time: _____
Online Time: _____

Type of Search
NA# _____ AA# _____
S/L: _____ Oligomer: _____
Encode/Transl: _____
Structure #: _____ Text: _____
Inventor: _____ Litigation: _____

Vendors and cost where applicable
STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other (Specify): _____

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OM protein - protein search, using sw model

Run on: January 25, 2006, 17:25:09 ; Search time 10 Seconds

(without alignments)
384.432 Million cell updates/sec

Title: US-10-767-521-1

Perfect score: 1854

Sequence: 1 MTTSDIVETFGTTSYDDV.....LERTSVSPSTAEPLSLIVF 355

Scoring table:

BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 75621 seqs, 10829074 residues

Total number of hits satisfying chosen parameters: 75621

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Published Applications AA New:*
1: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
4: /cgn2_6/ptodata/2/pubpaa/FCI_NEW_PUB.pep:*
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6: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
7: /cgn2_6/ptodata/2/pubpaa/US11_NEW_PUB.pep:*
8: /cgn2_6/ptodata/2/pubpaa/US60_NEW_PUB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1854	100.0	355	7	US-11-068-686-4
2	1854	100.0	355	7	US-11-127-877-64
3	1854	100.0	355	7	US-11-216-610-4
4	1846	99.6	355	7	US-11-216-610-2
5	1781	96.1	355	7	US-11-216-610-6
6	943.5	50.9	352	7	US-11-068-686-20
7	938.5	50.6	352	6	US-10-995-561-523
8	938.5	50.6	352	7	US-11-068-686-2
9	938.5	50.6	352	7	US-11-127-877-61
10	886.5	47.8	374	7	US-11-127-877-60
11	803.5	43.3	360	6	US-10-959-310-36
12	684	36.9	355	6	US-10-995-561-636
13	684	36.9	362	6	US-10-995-561-637
14	662.5	35.7	344	6	US-10-995-561-524
15	662.5	35.7	344	6	US-10-995-561-525
16	572	30.9	374	7	US-11-127-877-62
17	525.5	28.3	351	7	US-11-122-849-2
18	521	28.1	333	7	US-11-127-877-57
19	508	27.4	352	7	US-11-028-922A-1
20	506.5	27.3	216	6	US-10-995-561-522
21	503	27.1	353	7	US-11-017-058-9
22	492.5	26.6	415	7	US-11-017-058-2
23	480	25.3	349	7	US-11-028-922A-2
24	468.5	25.3	359	6	US-10-876-787-2
25	467	25.2	388	6	US-10-995-561-713

26	467	25.2	394	6	US-10-995-561-714	Sequence 714, App
27	467	25.2	394	6	US-10-995-561-715	Sequence 715, App
28	465.5	25.1	359	6	US-10-995-561-712	Sequence 712, App
29	465.5	25.1	359	6	US-10-995-561-716	Sequence 716, App
30	465.5	25.1	359	7	US-11-127-877-65	Sequence 65, App
31	462.5	24.9	254	6	US-10-055-877-248	Sequence 248, App
32	462.5	24.9	254	6	US-10-055-877-327	Sequence 327, App
33	462.5	24.9	254	6	US-10-055-877-340	Sequence 340, App
34	462.5	24.9	254	6	US-10-877-346-83	Sequence 83, App
35	445	24.0	229	6	US-10-055-877-225	Sequence 225, App
36	445	24.0	229	6	US-10-055-877-237	Sequence 237, App
37	417	22.5	391	7	US-11-127-877-52	Sequence 52, App
38	411.5	22.2	400	7	US-11-127-877-55	Sequence 55, App
39	382.5	20.6	388	6	US-10-995-561-838	Sequence 838, App
40	382.5	20.6	389	6	US-10-995-561-837	Sequence 837, App
41	382	20.6	269	7	US-11-151-482-5	Sequence 5, App
42	366.5	19.8	346	7	US-11-157-930-2	Sequence 2, App
43	362	19.5	339	7	US-11-157-930-4	Sequence 4, App
44	362	19.5	367	7	US-11-157-930-6	Sequence 6, App
45	354	19.1	350	6	US-10-502-145-1	Sequence 1, App

ALIGNMENTS

RESULT 1
US-11-068-686-4
Sequence 4, Application US/11068686
Publication No. US20050260565A1
GENERAL INFORMATION:
APPLICANT: Gray, Patrick W.
Schweickart, Vicki L.
Report, Carol J.
TITLE OF INVENTION: Chemokine Receptor Materials and Methods
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 S. Wacker Drive
CITY: Chicago
STATE: Illinois
COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/068,686
FILING DATE: 28-Feb-2005
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Noland, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 27866/33670
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
TELEFAX: 312-474-0448
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: /= "88-2b amino acid sequence"
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-11-068-686-4
Query Match 100.0%; Score 1854; DB 7; Length 355;
Best Local Similarity 100.0%; Pred. No. 3.4e-147;
Matches 355; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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QY      1 MTTSLDVTVEFTGTSYYDDVGLCEKADTRALMAQFVPLYSIVFTVGLGNVVMVMI 60
DB      1 MTTSLDVTVEFTGTSYYDDVGLCEKADTRALMAQFVPLYSIVFTVGLGNVVMVMI 60
QY      61 KYRLRIMTNIYILNLAISDLFLVTLPMIHYVRGHNWFGHGCKLLSGFYHTGLYSE 120
DB      61 KYRLRIMTNIYILNLAISDLFLVTLPMIHYVRGHNWFGHGCKLLSGFYHTGLYSE 120
QY      121 IFFIILITDRYLAIVHAVALRARTVGTITSIVTGLAVLAALPEFIETEBELFEE 180
DB      121 IFFIILITDRYLAIVHAVALRARTVGTITSIVTGLAVLAALPEFIETEBELFEE 180
QY      181 TCSALYPEDTVYSWNRHFTLMTIFCLVPLVNAICVTGIIKTLRCPSSKKYKARL 240
DB      181 TCSALYPEDTVYSWNRHFTLMTIFCLVPLVNAICVTGIIKTLRCPSSKKYKARL 240
QY      241 IFVIMAVFPIFWTPYVVAIILSSYOSILFGNDCERSKHLDMVLTEVIAYSHCCMNPVI 300
DB      241 IFVIMAVFPIFWTPYVVAIILSSYOSILFGNDCERSKHLDMVLTEVIAYSHCCMNPVI 300
QY      301 YAFVGERFRKRYLRHFFRHLMLHGRYIPFLPSEKLERISSVSPSTABPELSIVF 355
DB      301 YAFVGERFRKRYLRHFFRHLMLHGRYIPFLPSEKLERISSVSPSTABPELSIVF 355

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RESULT 2

```

US-11-127-877-64
; Sequence 64, Application US/11127877
; Publication No. US20050287565A1
; GENERAL INFORMATION:
; APPLICANT: Merchiers, Pascal G.
; APPLICANT: Hoffmann, Marcel
; APPLICANT: Spittaels, Koensraad F. F.
; APPLICANT: Laenen, Wendy
; TITLE OF INVENTION: Method, Compositions and Compound Assays For Inhibiting
; TITLE OF INVENTION: Amyloid-beta Protein Production
; FILE REFERENCE: P27,800-B USA
; CURRENT APPLICATION NUMBER: US/11/127,877
; PRIOR FILING DATE: 2005-05-12
; PRIOR APPLICATION NUMBER: 60/570,352
; PRIOR FILING DATE: 2004-05-12
; PRIOR APPLICATION NUMBER: 60/603,948
; PRIOR FILING DATE: 2004-08-24
; NUMBER OF SEQ ID NOS: 590
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 64
; LENGTH: 355
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-127-877-64

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Query Match      100.0%; Score 1854; DB 7; Length 355;
Best Local Similarity 100.0%; Pred. No. 3,4e-147;
Matches 355; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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QY      1 MTTSLDVTVEFTGTSYYDDVGLCEKADTRALMAQFVPLYSIVFTVGLGNVVMVMI 60
DB      1 MTTSLDVTVEFTGTSYYDDVGLCEKADTRALMAQFVPLYSIVFTVGLGNVVMVMI 60
QY      61 KYRLRIMTNIYILNLAISDLFLVTLPMIHYVRGHNWFGHGCKLLSGFYHTGLYSE 120
DB      61 KYRLRIMTNIYILNLAISDLFLVTLPMIHYVRGHNWFGHGCKLLSGFYHTGLYSE 120
QY      121 IFFIILITDRYLAIVHAVALRARTVGTITSIVTGLAVLAALPEFIETEBELFEE 180
DB      121 IFFIILITDRYLAIVHAVALRARTVGTITSIVTGLAVLAALPEFIETEBELFEE 180
QY      181 TCSALYPEDTVYSWNRHFTLMTIFCLVPLVNAICVTGIIKTLRCPSSKKYKARL 240
DB      181 TCSALYPEDTVYSWNRHFTLMTIFCLVPLVNAICVTGIIKTLRCPSSKKYKARL 240
QY      241 IFVIMAVFPIFWTPYVVAIILSSYOSILFGNDCERSKHLDMVLTEVIAYSHCCMNPVI 300
DB      241 IFVIMAVFPIFWTPYVVAIILSSYOSILFGNDCERSKHLDMVLTEVIAYSHCCMNPVI 300

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DB      241 IFVIMAVFPIFWTPYVVAIILSSYOSILFGNDCERSKHLDMVLTEVIAYSHCCMNPVI 300
QY      301 YAFVGERFRKRYLRHFFRHLMLHGRYIPFLPSEKLERISSVSPSTABPELSIVF 355
DB      301 YAFVGERFRKRYLRHFFRHLMLHGRYIPFLPSEKLERISSVSPSTABPELSIVF 355

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RESULT 3

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US-11-216-610-4
; Sequence 4, Application US/11216610
; Publication No. US20060002926A1
; GENERAL INFORMATION:
; APPLICANT: Gerard, Craig J.
; APPLICANT: Gerard, Norma P.
; APPLICANT: Mackay, Charles R.
; APPLICANT: Ponath, Paul D.
; APPLICANT: Post, Theodore W.
; APPLICANT: Qin, Shixin
; TITLE OF INVENTION: G PROTEIN-COUPLED RECEPTOR GENE CCR3 AND
; ANTAGONISTS THEREOF
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
; STREET: Two Milltia Drive
; CITY: Lexington
; STATE: MA
; COUNTRY: USA
; ZIP: 02173

```

COMPUTER READABLE FORM:

```

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

```

CURRENT APPLICATION DATA:

```

APPLICATION NUMBER: US/11/216,610
FILING DATE: 31-Aug-2005
CLASSIFICATION: 536

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PRIOR APPLICATION DATA:

```

APPLICATION NUMBER: US/08/963,656
FILING DATE: 03-NOV-1997
APPLICATION NUMBER: 08/720,565
FILING DATE: 30-SEP-1996
APPLICATION NUMBER: PCT/US96/00608
FILING DATE: 19-JAN-1996
APPLICATION NUMBER: US 08/375,199
FILING DATE: 19-JAN-1995

```

ATTORNEY/AGENT INFORMATION:

```

NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: LKS94-05A22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
TELEFAX: 781-861-9540

```

INFORMATION FOR SEQ ID NO: 4:

```

SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
TYPE: amino acid
MOLECULE TYPE: protein
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-11-216-610-4

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Query Match      100.0%; Score 1854; DB 7; Length 355;
Best Local Similarity 100.0%; Pred. No. 3,4e-147;
Matches 355; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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QY      1 MTTSLDVTVEFTGTSYYDDVGLCEKADTRALMAQFVPLYSIVFTVGLGNVVMVMI 60
DB      1 MTTSLDVTVEFTGTSYYDDVGLCEKADTRALMAQFVPLYSIVFTVGLGNVVMVMI 60
QY      61 KYRLRIMTNIYILNLAISDLFLVTLPMIHYVRGHNWFGHGCKLLSGFYHTGLYSE 120
DB      61 KYRLRIMTNIYILNLAISDLFLVTLPMIHYVRGHNWFGHGCKLLSGFYHTGLYSE 120

```

MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 2:

NAME: BLOOM, DAVID B.
REGISTRATION NUMBER: 22,592

REFERENCE/DOCKET NUMBER: LKS94-05A22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-11-216-610-6

Query Match 96.1%; Score 1781; DB 7; Length 355;
Best Local Similarity 96.6%; Pred. No. 3.9e-141;
Matches 343; Conservative 0; Mismatches 12; Indels 0; Gaps 0;

QY 1 MTTSDVETFGTGYDDVGLCEKADTRALMAQFVPPYLSLVFTVGLGNVVMVILI 60
DB 1 MTTSDVETFGTGYDDVGLCEKADTRALMAQFVPPYLSLVFTVGLGNVVMVILI 60
QY 61 KYRLRLIMTNYILMLAISDLFLVTLFPWIIHYVGHNVFSGHCKLISGFYHTGLYSE 120
DB 61 KYRLRLIMTNYILMLAISDLFLVTLFPWIIHYVGHNVFSGHCKLISGFYHTGLYSE 120
QY 121 IFFIILITDRYLAIVHAFALRARTVFGVTSIVTWGLAVIALPFIETETELPERE 180
DB 121 IFFIILITDRYLAIVHAFALRARTVFGVTSIVTWGLAVIALPFIETETELPERE 180
QY 181 TLCSALYPEDVYVSRHFFTLMTIFCLVLPILVMAICYTGIIKTLRCPSKKKYAIKL 240
DB 181 TYCSALYPEDVYVSRHFFTLMTIFCLVLPILVMAICYTGIIKTLRCPSKKKYAIKL 240
QY 241 IFVIMAVFPIFWTPYVVAIILSXXKXILFGNDCERXXXXLVMLVTEVIAVSHCCNPIY 300
DB 241 IFVIMAVFPIFWTPYVVAIILSXXKXILFGNDCERXXXXLVMLVTEVIAVSHCCNPIY 300
QY 301 YAFVGRFRPKYLRFPHRLMLHGLGYIFLPSEKLERTSSVSPSTAEPELSIVF 355
DB 301 YAFVGRFRPKYLRFPHRLMLHGLGYIFLPSEKLERTSSVSPSTAEPELSIVF 355

RESULT 6
US-11-068-686-20
Sequence 20, Application US/11068686
Publication No. US20050260565A1
GENERAL INFORMATION:
APPLICANT: Gray, Patrick W.
Schweickart, Vicky L.
Report, Carol J.
TITLE OF INVENTION: Chemokine Receptor Materials and Methods
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 S. Wacker Drive
CITY: Chicago
STATE: Illinois
COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent in Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/068,686
FILING DATE: 28-Feb-2005
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Noland, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 27866/33670

TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
TELEFAX: 312-474-0448
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 352 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 20:
US-11-068-686-20

Query Match 50.9%; Score 943.5; DB 7; Length 352;
Best Local Similarity 54.2%; Pred. No. 1.4e-71;
Matches 180; Conservative 58; Mismatches 91; Indels 3; Gaps 3;

QY 24 CEKADTRALMAQFVPPYLSLVFTVGLGNVVMVILIKYRLRLIMTNYILMLAISDLFL 83
DB 20 CQKINVQIAARLLPPLYSLVIFGFGNMVILILINCKRLKSMTDIYILMLAISDLFL 79
QY 84 LVTLFPWIIHYVGHNVFSGHCKLISGFYHTGLYSEIFFIILITDRYLAIVHAFALR 143
DB 80 LITVFPMAHYAAA-QMDGNTMCQULTGLYIFGFSGIFFIILITDRYLAIVHAFALR 138
QY 144 ARTVFGVTSIVTWGLAVIALPFIETETELPEETLCSALYPEDVYVSRHFFHILRM 203
DB 139 ARTVFGVTSIVTWGLAVIALPFIETETELPEETLCSALYPEDVYVSRHFFHILRM 198
QY 204 TIFCLVPLVMAICYTGIIKTLRCPK-KKKYARILIFVIMAVFPIFWTPYVVAIILS 262
DB 199 VILGLVPLVMAICYSGILKTLKCRNEKRRHRAVRLFTIMVYFLMLAPYVILNLN 258
QY 263 SYGSLIFGDCERSKHLVLMVTEVIAVSHCCNPIYAFVGERFRKYLRFHFRHLLM 322
DB 259 TPQEFGLANCGSSRLDQAMQVTTLTGHTCCINPIIYAFVGERFRYLVLFQKHLAK 318
QY 323 HLGRIYIPLPSEKLERTSSV-SPSTAEPELSI 353
DB 319 RFCKCISIFQGEAPERASSVYTRSTGEQELISV 350

RESULT 7
US-10-995-561-523
Sequence 523, Application US/10995561
Publication No. US20050272054A1
GENERAL INFORMATION:
APPLICANT: Cargill, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: C1001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 523
LENGTH: 352
TYPE: PRT
ORGANISM: Homo sapiens
US-10-995-561-523

Query Match 50.6%; Score 938.5; DB 6; Length 352;
Best Local Similarity 53.3%; Pred. No. 3.5e-71;
Matches 177; Conservative 62; Mismatches 90; Indels 3; Gaps 3;

QY 24 CEKADTRALMAQFVPPYLSLVFTVGLGNVVMVILIKYRLRLIMTNYILMLAISDLFL 83
DB 20 CQKINVQIAARLLPPLYSLVIFGFGNMVILILINCKRLKSMTDIYILMLAISDLFL 79
QY 84 LVTLFPWIIHYVGHNVFSGHCKLISGFYHTGLYSEIFFIILITDRYLAIVHAFALR 143
DB 80 LITVFPMAHYAAA-QMDFGNTMCQULTGLYIFGFSGIFFIILITDRYLAIVHAFALR 138

